

<110> Jager, Dirk
Stockert, Elizabeth
Scanlan, Matthew
Gure, Ali
Knuth, Alexander
Old, Lloyd
Chen, Yao-tseng

<120> Isolated Nucleic Acid Molecules Encoding Cancer Associated Antigens,
the Antigens Per Se, and Uses Thereof

<130> LUD 5793.1

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<150> US 09/451,739

<151> 2000-06-22

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<151> 1999-11-30

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<212> DNA

<213> Homo sapiens

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<221> CDS

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cccatcgacc	ccaacgaacc	cacgtactgt	ctgtgcaacc	aggtctccta	tggggagatg	660
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<211> 857

<212> DNA

<213> Homo sapiens

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20          25          30
Asn Val Ser Leu Met Arg Glu Ile Asp Ala Lys Tyr Gln Glu Ile Leu
35          40          45
Lys Glu Leu Asp Glu Cys Tyr Glu Arg Phe Ser Arg Glu Thr Asp Gly
50          55          60
Ala Gln Lys Arg Arg Met Leu His Cys Val Gln Arg Ala Leu Ile Arg
65          70          75          80
Ser Gln Glu Leu Gly Asp Glu Lys Ile Gln Ile Val Ser Gln Met Val
85          90          95
Glu Leu Val Glu Asn Arg Thr Arg Gln Val Asp Ser His Val Glu Leu
100         105         110
Phe Glu Ala Gln Gln Glu Leu Gly Asp Thr Val Gly Asn Ser Gly Lys
115         120         125
Val Gly Ala Asp Arg Pro Asn Gly Asp Ala Val Ala Gln Ser Asp Lys
130         135         140
Pro Asn Ser Lys Arg Ser Arg Arg Gln Arg Asn Asn Glu Asn Arg Glu
145         150         155         160
Asn Ala Ser Ser Asn His Asp His Asp Asp Gly Ala Ser Gly Thr Pro
165         170         175
Lys Glu Lys Lys Ala Lys Thr Ser Lys Lys Lys Arg Ser Lys Ala
180         185         190
Lys Ala Glu Arg Glu Ala Ser Pro Ala Asp Leu Pro Ile Asp Pro Asn
195         200         205
Glu Pro Thr Tyr Cys Leu Cys Asn Gln Val Ser Tyr Gly Glu Met Ile
210         215         220
Gly Cys Asp Asn Asp Glu Cys Pro Ile Glu Trp Phe His Phe Ser Cys
225         230         235         240
Val Gly Leu Asn His Lys Pro Lys Gly Lys Trp Tyr Cys Pro Lys Cys
245         250         255
Arg Gly Glu Asn Glu Lys Thr Met Asp Lys Ala Leu Glu Lys Ser Lys
260         265         270
Lys Glu Arg Ala Tyr Asn Arg
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20        25        30
Arg Thr Arg Gln Val Asp Ser His Val Glu Leu Phe Glu Ala Gln Gln
35        40        45
Glu Leu Gly Asp Thr Val Gly Asn Ser Gly Lys Val Gly Ala Asp Arg
50        55        60
Pro Asn Gly Asp Ala Val Ala Gln Ser Asp Lys Pro Asn Ser Lys Arg
65        70        75        80
Ser Arg Arg Gln Arg Asn Asn Glu Asn Arg Glu Asn Ala Ser Ser Asn
85        90        95
His Asp His Asp Asp Gly Ala Ser Gly Thr Pro Lys Glu Lys Lys Ala
100       105       110
Lys Thr Ser Lys Lys Lys Lys Arg Ser Lys Ala Lys Ala Glu Arg Glu
115       120       125
Ala Ser Pro Ala Asp Leu Pro Ile Asp Pro Asn Glu Pro Thr Tyr Cys
130       135       140
Leu Cys Asn Gln Val Ser Tyr Gly Glu Met Ile Gly Cys Asp Asn Asp
145       150       155       160
Glu Cys Pro Ile Glu Trp Phe His Phe Ser Cys Val Gly Leu Asn His
165       170       175
Lys Pro Lys Gly Lys Trp Tyr Cys Pro Lys Cys Arg Gly Glu Asn Glu
180       185       190
Lys Thr Met Asp Lys Ala Leu Glu Lys Ser Lys Lys Glu Arg Ala Tyr
195       200       205
Asn Arg
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<210> 7
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<213> Homo sapiens
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20        25        30
Ala Leu Ile Arg Ser Gln Glu Leu Gly Asp Glu Lys Ile Gln Ile Val
35        40        45
Ser Gln Met Val Glu Leu Val Glu Asn Arg Thr Arg Gln Val Asp Ser
50        55        60
His Val Glu Leu Phe Glu Ala Gln Gln Glu Leu Gly Asp Thr Val Gly
65        70        75        80
Asn Ser Gly Lys Val Gly Ala Asp Arg Pro Asn Gly Asp Ala Val Ala
85        90        95
Gln Ser Asp Lys Pro Asn Ser Lys Arg Ser Arg Arg Gln Arg Asn Asn
100       105       110
Glu Asn Arg Glu Asn Ala Ser Ser Asn His Asp His Asp Asp Gly Ala
115       120       125

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Ser Gly Thr Pro Lys Glu Lys Lys Ala Lys Thr Ser Lys Lys Lys Lys
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 Arg Ser Lys Ala Lys Ala Glu Arg Glu Ala Ser Pro Ala Asp Leu Pro
 145 150 155 160
 Ile Asp Pro Asn Glu Pro Thr Tyr Cys Leu Cys Asn Gln Val Ser Tyr
 165 170 175
 Gly Glu Met Ile Gly Cys Asp Asn Asp Glu Cys Pro Ile Glu Trp Phe
 180 185 190
 His Phe Ser Cys Val Gly Leu Asn His Lys Pro Lys Gly Lys Trp Tyr
 195 200 205
 Cys Pro Lys Cys Arg Gly Glu Asn Glu Lys Thr Met Asp Lys Ala Leu
 210 215 220
 Glu Lys Ser Lys Lys Glu Arg Ala Tyr Asn Arg
 225 230 235

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<210> 11
 <211> 21
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<213> Homo sapiens
 <400> 11
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<210> 12
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 <213> Homo sapiens
 <400> 12
 cgtggtcgtg gttgctggac gcg 23

<210> 13
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<210> 14
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 cgtggtcgtg gttgctggac gcg 23

<210> 15
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<210> 16
<211> 513
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Ile Glu Met Gln Lys Ser Val Pro Asn Lys Ala Leu Glu Leu Lys Asn
35        40        45
Glu Gln Thr Leu Arg Ala Asp Glu Ile Leu Pro Ser Glu Ser Lys Gln
50        55        60
Lys Asp Tyr Glu Glu Ser Ser Trp Asp Ser Glu Ser Leu Cys Glu Thr
65        70        75        80
Val Ser Gln Lys Asp Val Cys Leu Pro Lys Ala Thr His Gln Lys Glu
85        90        95
Ile Asp Lys Ile Asn Gly Lys Leu Glu Glu Ser Pro Asp Asn Asp Gly
100       105       110
Phe Leu Lys Ala Pro Cys Arg Met Lys Val Ser Ile Pro Thr Lys Ala
115       120       125
Leu Glu Leu Met Asp Met Gln Thr Phe Lys Ala Glu Pro Pro Glu Lys
130       135       140
Pro Ser Ala Phe Glu Pro Ala Ile Glu Met Gln Lys Ser Val Pro Asn
145       150       155       160
Lys Ala Leu Glu Leu Lys Asn Glu Gln Thr Leu Arg Ala Asp Gln Met
165       170       175
Phe Pro Ser Glu Ser Lys Gln Lys Lys Val Glu Glu Asn Ser Trp Asp
180       185       190
Ser Glu Ser Leu Arg Glu Thr Val Ser Gln Lys Asp Val Cys Val Pro
195       200       205
Lys Ala Thr His Gln Lys Glu Met Asp Lys Ile Ser Gly Lys Leu Glu
210       215       220
Asp Ser Thr Ser Leu Ser Lys Ile Leu Asp Thr Val His Ser Cys Glu
225       230       235       240
Arg Ala Arg Glu Leu Gln Lys Asp His Cys Glu Gln Arg Thr Gly Lys
245       250       255
Met Glu Gln Met Lys Lys Lys Phe Cys Val Leu Lys Lys Lys Leu Ser
260       265       270
Glu Ala Lys Glu Ile Lys Ser Gln Leu Glu Asn Gln Lys Val Lys Trp
275       280       285
Glu Gln Glu Leu Cys Ser Val Arg Leu Thr Leu Asn Gln Glu Glu Glu

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290	295	300
Lys Arg Arg Asn Ala Asp	Ile Leu Asn Glu Lys	Ile Arg Glu Glu Leu
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Gly Arg Ile Glu Glu Gln	His Arg Lys Glu Leu	Glu Val Lys Gln Gln
	325	330
Leu Glu Gln Ala Leu Arg	Ile Gln Asp Ile Glu	Leu Lys Ser Val Glu
	340	345
Ser Asn Leu Asn Gln Val	Ser His Thr His Glu	Asn Glu Asn Tyr Leu
	350	355
Leu His Glu Asn Cys Met	Leu Lys Lys Glu Ile	Ala Met Leu Lys Leu
	360	365
Glu Ile Ala Thr Leu Lys	His Gln Tyr Gln Glu	Lys Glu Asn Lys Tyr
370	375	380
Phe Glu Asp Ile Lys Ile	Leu Lys Glu Lys Asn	Ala Glu Leu Gln Met
	385	390
Thr Leu Lys Leu Lys Glu	Glu Ser Leu Thr Lys	Arg Ala Ser Gln Tyr
	395	400
Ser Gly Gln Leu Lys Val	Leu Ile Ala Glu Asn	Thr Met Leu Thr Ser
	405	410
Lys Leu Lys Glu Lys Gln	Asp Lys Glu Ile Leu	Glu Ala Glu Ile Glu
	415	420
Ser His His Pro Arg Leu	Ala Ser Ala Val Gln	Asp His Asp Gln Ile
	425	430
Val Thr Ser Arg Lys Ser	Gln Glu Pro Ala Phe	His Ile Ala Gly Asp
	435	440
Ala Cys Leu Gln Arg Lys	Met Asn Val Asp Val	Ser Ser Thr Asp Ile
	445	450
	455	460
	465	470
	475	480
	485	490
	495	500
	505	510

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<210> 19
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 Asp Pro Gly Pro Val Ala Arg Gly Arg Gly Cys Ser Ser Asp Arg Leu
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 Pro Arg Pro Ala Gly Pro Ala Arg Arg Gln Phe Gln Ala Ala Ser Leu
 35 40 45
 Leu Thr Arg Gly Trp Gly Arg Ala Trp Pro Trp Lys Gln Ile Leu Lys

50	55	60
Glu Leu Asp Glu Cys Tyr Glu Arg Phe Ser Arg Glu Thr Asp Gly Ala		
65	70	75
Gln Lys Arg Arg Met Leu His Cys Val Gln Arg Ala Leu Ile Arg Ser		80
	85	90
Gln Glu Leu Gly Asp Glu Lys Ile Gln Ile Val Ser Gln Met Val Glu		95
	100	105
Leu Val Glu Asn Arg Thr Arg Gln Val Asp Ser His Val Glu Leu Phe		110
	115	120
Glu Ala Gln Gln Glu Leu Gly Asp Thr Val Gly Asn Ser Gly Lys Val		125
	130	135
Gly Ala Asp Arg Pro Asn Gly Asp Ala Val Ala Gln Ser Asp Lys Pro		140
145	150	155
Asn Ser Lys Arg Ser Arg Arg Gln Arg Asn Asn Glu Asn Arg Glu Asn		160
	165	170
Ala Ser Ser Asn His Asp His Asp Asp Gly Ala Ser Gly Thr Pro Lys		175
	180	185
Glu Lys Lys Ala Lys Thr Ser Lys Lys Lys Lys Arg Ser Lys Ala Lys		190
	195	200
Ala Glu Arg Glu Ala Ser Pro Ala Asp Leu Pro Ile Asp Pro Asn Glu		205
	210	215
Pro Thr Tyr Cys Leu Cys Asn Gln Val Ser Tyr Gly Glu Met Ile Gly		220
225	230	235
Cys Asp Asn Asp Glu Cys Pro Ile Glu Trp Phe His Phe Ser Cys Val		240
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Ser Val Val Ala Lys Leu Leu Ser His Gly Ala Val Ile Glu Val His
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225         230         235         240
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Thr Pro Asp Thr Ala Glu Ser Leu Val Glu Lys Thr Pro Asp Glu Ala
          260         265         270
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25362054.1

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Pro	Pro	Glu	Lys	Pro	Ser	Ala	Phe	Glu	Pro	Ala	Ile	Glu	Met	Gln	Lys		820	825	830
Ser	Val	Pro	Asn	Lys	Ala	Leu	Glu	Leu	Lys	Asn	Glu	Gln	Thr	Leu	Arg		835	840	845
Ala	Asp	Gln	Met	Phe	Pro	Ser	Glu	Ser	Lys	Gln	Lys	Lys	Val	Glu	Glu		850	855	860
Asn	Ser	Trp	Asp	Ser	Glu	Ser	Leu	Arg	Glu	Thr	Val	Ser	Gln	Lys	Asp	865	870	875	880
Val	Cys	Val	Pro	Lys	Ala	Thr	His	Gln	Lys	Glu	Met	Asp	Lys	Ile	Ser		885	890	895
Gly	Lys	Leu	Glu	Asp	Ser	Thr	Ser	Leu	Ser	Lys	Ile	Leu	Asp	Thr	Val		900	905	910
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Ser	Thr	Ile	Tyr	Asn	Asn	Glu	Val	Leu	His	Gln	Pro	Leu	Ser	Glu	Ala	1185	1190	1195	1200
Gln	Arg	Lys	Ser	Lys	Ser	Leu	Lys	Ile	Asn	Leu	Asn	Tyr	Ala	Gly	Asp		1205	1210	1215
Ala	Leu	Arg	Glu	Asn	Thr	Leu	Val	Ser	Glu	His	Ala	Gln	Arg	Asp	Gln		1220	1225	1230
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Gln Lys Leu Phe Gln Leu Gln Ser Lys Asn Met Trp Leu Gln Gln Gln		
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Lys Asn Glu Glu Ile Phe Asn Tyr Asn Asn His Leu Lys Asn Arg Ile		
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1060 1065 1070
Leu Arg Ile Gln Asp Ile Glu Leu Lys Ser Val Glu Ser Asn Leu Asn
1075 1080 1085
Gln Val Ser His Thr His Glu Asn Glu Asn Tyr Leu Leu His Glu Asn
1090 1095 1100
Cys Met Leu Lys Lys Glu Ile Ala Met Leu Lys Leu Glu Ile Ala Thr
1105 1110 1115 1120
Leu Lys His Gln Tyr Gln Glu Lys Glu Asn Lys Tyr Phe Glu Asp Ile
1125 1130 1135
Lys Ile Leu Lys Glu Lys Asn Ala Glu Leu Gln Met Thr Leu Lys Leu
1140 1145 1150
Lys Glu Glu Ser Leu Thr Lys Arg Ala Ser Gln Tyr Ser Gly Gln Leu
1155 1160 1165
Lys Val Leu Ile Ala Glu Asn Thr Met Leu Thr Ser Lys Leu Lys Glu
1170 1175 1180
Lys Gln Asp Lys Glu Ile Leu Glu Ala Glu Ile Glu Ser His His Pro
1185 1190 1195 1200
Arg Leu Ala Ser Ala Val Gln Asp His Asp Gln Ile Val Thr Ser Arg
1205 1210 1215
Lys Ser Gln Glu Pro Ala Phe His Ile Ala Gly Asp Ala Cys Leu Gln
1220 1225 1230
Arg Lys Met Asn Val Asp Val Ser Ser Thr Ile Tyr Asn Asn Glu Val
1235 1240 1245
Leu His Gln Pro Leu Ser Glu Ala Gln Arg Lys Ser Lys Ser Leu Lys
1250 1255 1260
Ile Asn Leu Asn Tyr Ala Gly Asp Ala Leu Arg Glu Asn Thr Leu Val
1265 1270 1275 1280
Ser Glu His Ala Gln Arg Asp Gln Arg Glu Thr Gln Cys Gln Met Lys
1285 1290 1295
Glu Ala Glu His Met Tyr Gln Asn Glu Gln Asp Asn Val Asn Lys His
1300 1305 1310
Thr Glu Gln Gln Glu Ser Leu Asp Gln Lys Leu Phe Gln Leu Gln Ser
1315 1320 1325
Lys Asn Met Trp Leu Gln Gln Gln Leu Val His Ala His Lys Lys Ala
1330 1335 1340
Asp Asn Lys Ser Lys Ile Thr Ile Asp Ile His Phe Leu Glu Arg Lys
1345 1350 1355 1360
Met Gln His His Leu Lys Glu Lys Asn Glu Glu Ile Phe Asn Tyr
1365 1370 1375
Asn Asn His Leu Lys Asn Arg Ile Tyr Gln Tyr Glu Lys Glu Lys Ala
1380 1385 1390
Glu Thr Glu Asn Ser
1395